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From terminology integration to information integration

An example in the domain of genetics





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Outline

- Background
 - Terminology integration: The Unified Medical Language System
 - Information integration: *Genetics as an example*
- Applications
 - GenesTrace
 - BioMeKe



Terminology integration

The Unified Medical Language System

Motivation

- ◆ Started in 1986
- National Library of Medicine
- "Long-term R&D project"
- Complementary to IAIMS

(Integrated Academic Information Management Systems)

- «[...] the UMLS project is an effort to overcome two significant barriers to effective retrieval of machine-readable information.
- The first is the variety of ways the same concepts are expressed in different machine-readable sources and by different people.
- The second is the distribution of useful information among many disparate databases and systems.»



Source Vocabularies

- **♦** 117 "sources"
- ◆ ~60 families of vocabularies
 - multiple translations (e.g., MeSH, ICPC, ICD-10)
 - variants (American-English equivalents, Australian extension/adaptation)
 - subsequent versions usually considered distinct families (ICD: 9-10; DSM: IIIR-IV)
- Broad coverage of biomedicine
- Common presentation



Biomedical terminologies

- ◆ Core vocabularies
 - anatomy (UWDA, Neuronames)
 - drugs (First DataBank, Micromedex)
 - medical devices (UMD, SPN)
- Several perspectives
 - clinical terms (SNOMED, CTV3)
 - information sciences (MeSH, CRISP)
 - administrative terminologies (ICD-9-CM, CPT-4)
 - standards (HL7, LOINC)

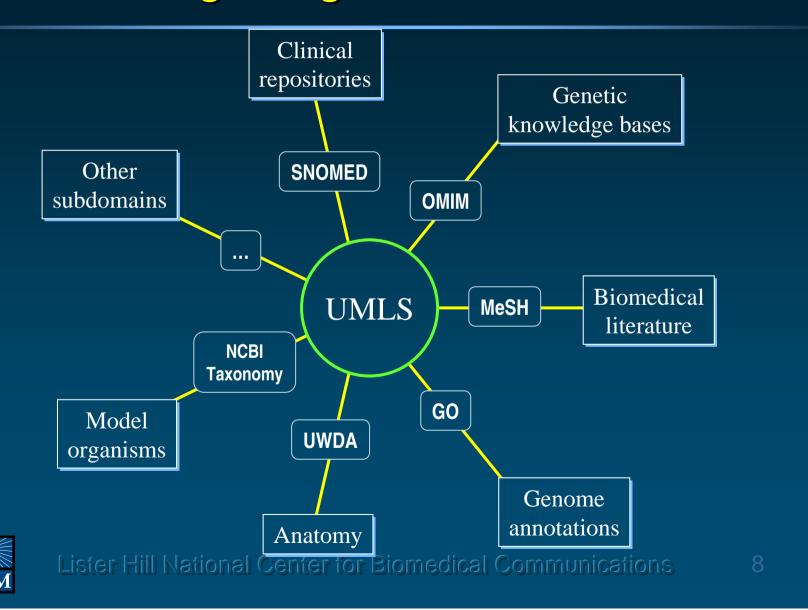


Biomedical terminologies (cont'd)

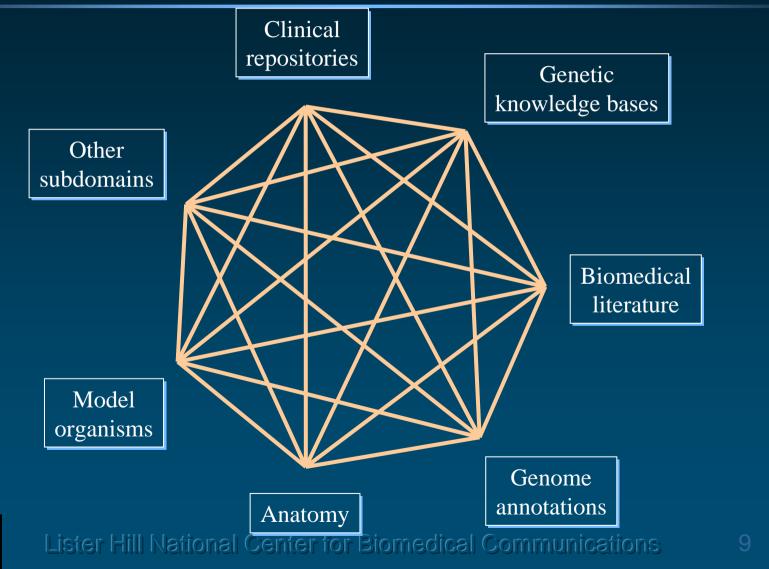
- Specialized vocabularies
 - nursing (NIC, NOC, NANDA, Omaha, PCDS)
 - dentistry (CDT)
 - oncology (PDQ)
 - psychiatry (DSM, APA)
 - adverse reactions (COSTART, WHO ART)
 - primary care (ICPC)
- ◆ Knowledge bases (AI/Rheum, DXplain, QMR)



Integrating subdomains



Integrating subdomains



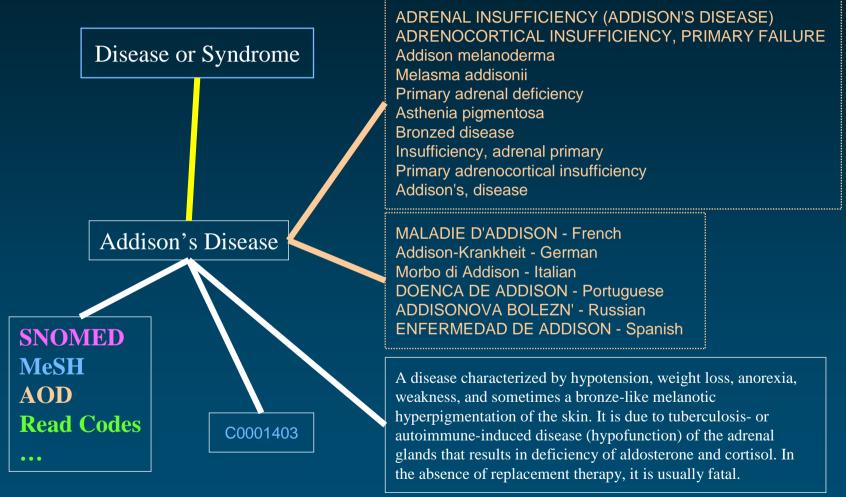


UMLS: 3 components

- ◆ Metathesaurus
 - Concepts
 - Inter-concept relationships
- ◆ Semantic Network
 - Semantic types
 - Semantic network relationships
- **♦** Lexical resources
 - SPECIALIST Lexicon
 - Lexical tools



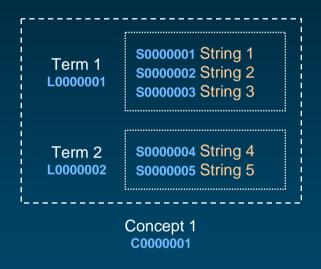
Addison's Disease: Concept





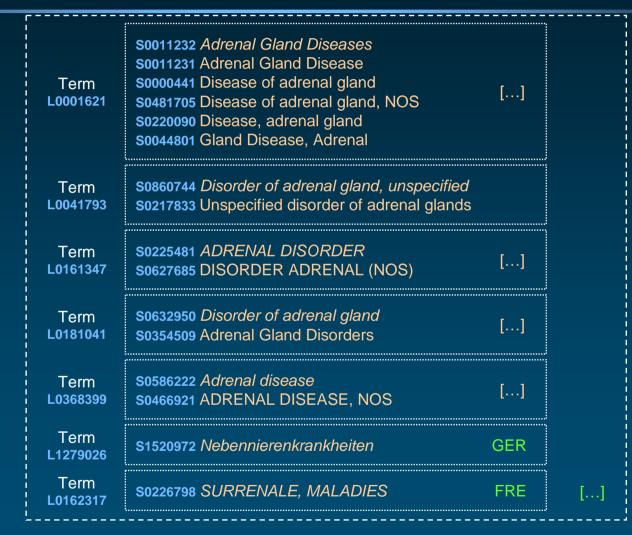
Metathesaurus Concepts (2003AA)

- Concept: Cluster of synonymous terms
 - ~875,000 concepts
 - identified by a CUI
- ◆ Term: Set of lexical variants
 - ~1.8 M terms
 - identified by a LUI
- ◆ String: Concept name
 - ~2.1 M strings
 - identified by a SUI





Cluster of synonymous terms





Concept

C0001621

Metathesaurus Relationships

◆ Symbolic relations: ~5 M pairs of concepts

◆ Statistical relations : ~6.5 M pairs of concepts (co-occurring concepts)

 Categorization: Relationships between concepts and semantic types from the Semantic Network



Symbolic relations

◆ Relation MRREL

- Pair of concept identifiers
- Type
- Attribute (if any)
- List of sources (for type and attribute)
- ◆ Semantics of the relationship: defined by its type [and attribute]



Symbolic relationships Type

◆ Hierarchical

PAR/CHD Parent / Child

RB/RN Broader / Narrower than



Derived from hierarchies

• Siblings (children of parents) SIB



Other RO



Various flavors of near-synonymy

Similar RL

 Source asserted synonymy SY

Possible synonymy RO

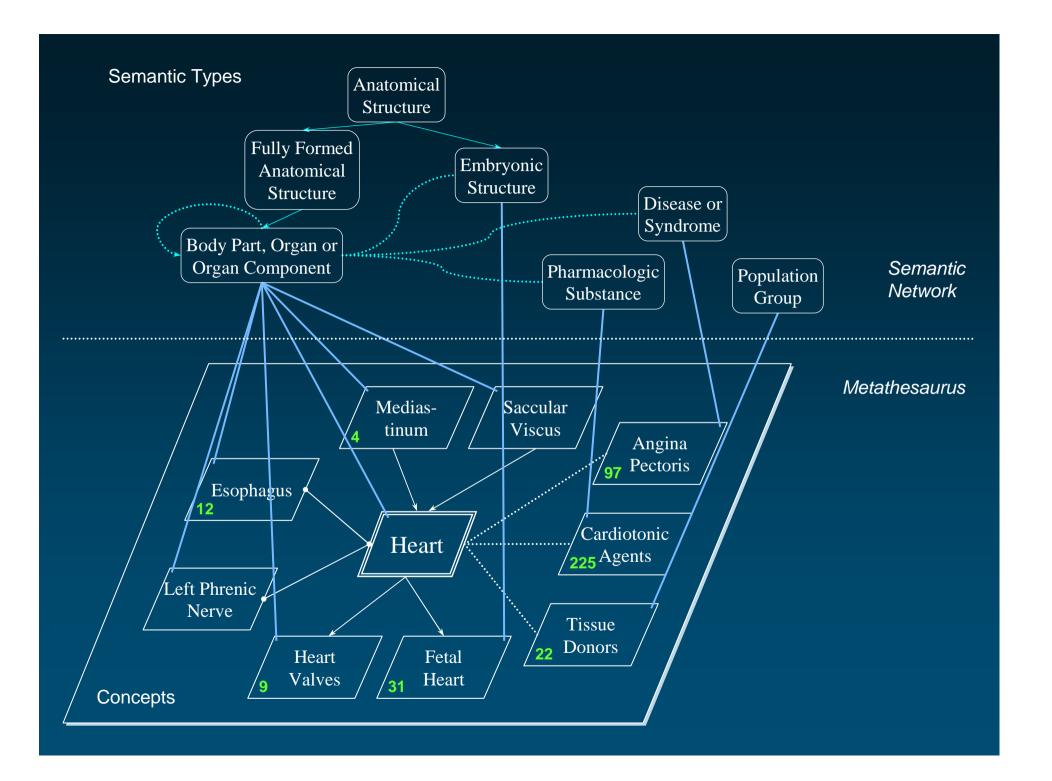




Symbolic relationships Attribute

- ◆ Hierarchical
 - isa (is-a-kind-of)
 - part-of
- **♦** Associative
 - location-of
 - caused-by
 - treats
 - ...
- Cross-references (mapping)



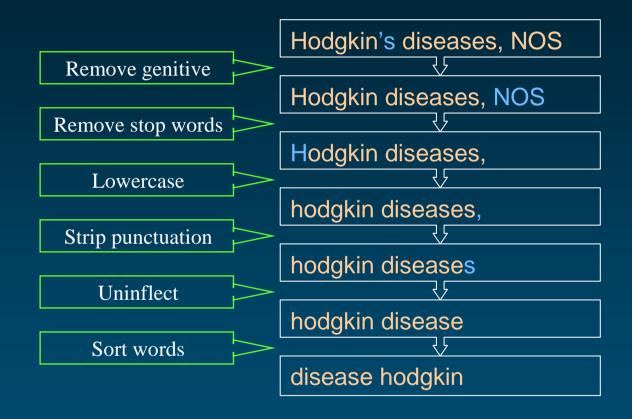


Lexical tools

- ◆ To manage lexical variation in biomedical terminologies
- Major tools
 - Normalization
 - Indexes
 - Lexical Variant Generation program (lvg)
- ◆ Based on the SPECIALIST Lexicon
- ◆ Used by noun phrase extractors, search engines



Normalization





Normalization: Example

Hodgkin Disease **HODGKINS DISEASE** Hodgkin's Disease Disease, Hodgkin's Hodgkin's, disease HODGKIN'S DISEASE Hodgkin's disease **Hodgkins Disease** Hodgkin's disease NOS Hodgkin's disease, NOS Disease, Hodgkins Diseases, Hodgkins **Hodgkins Diseases** Hodgkins disease hodgkin's disease Disease, Hodgkin

normalize disease hodgkin



Information integration

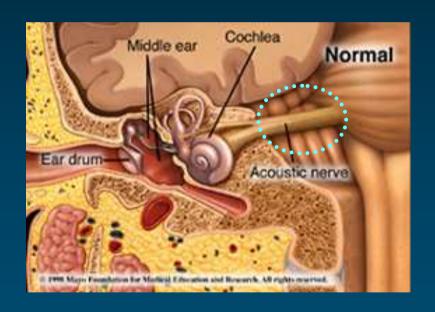
Genetics as an example

NF2 Gene, protein, and disease

Neurofibromatosis 2 is an autosomal dominant disease characterized by tumors called schwannomas involving the acoustic nerve, as well as other features. The disorder is caused by mutations of the NF2 gene resulting in absence or inactivation of the protein product. The protein product of NF2 is commonly called merlin (but also neurofibromin 2 and schwannomin) and functions as a tumor suppressor.



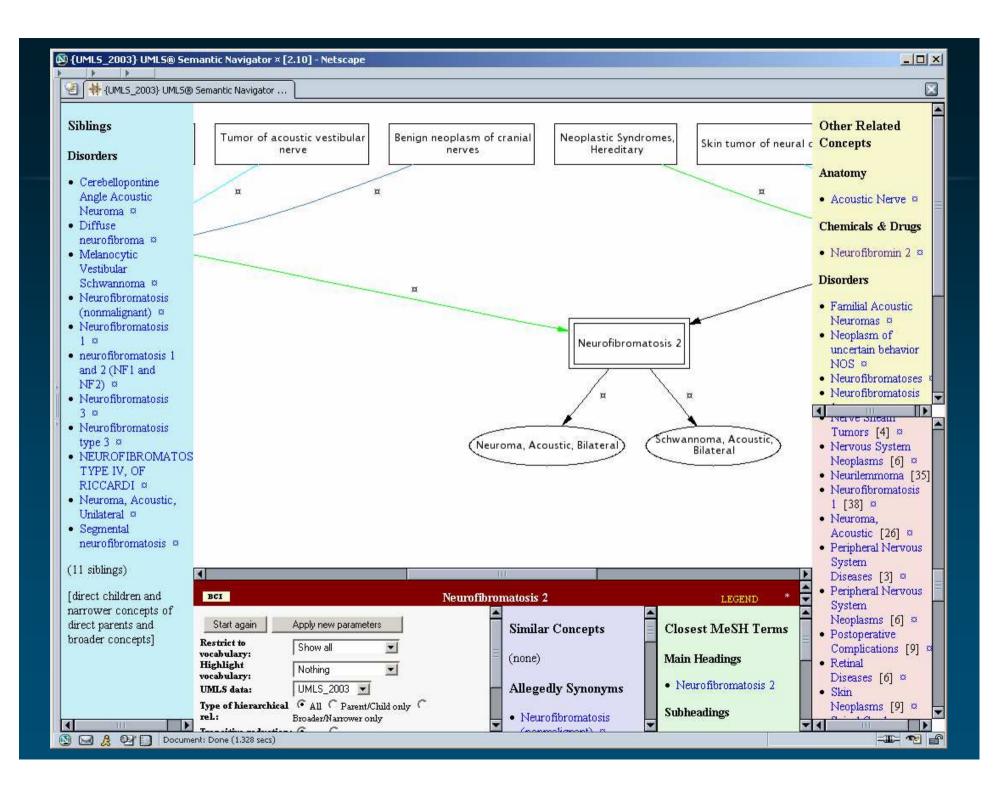
Schwannoma (acoustic neuroma)



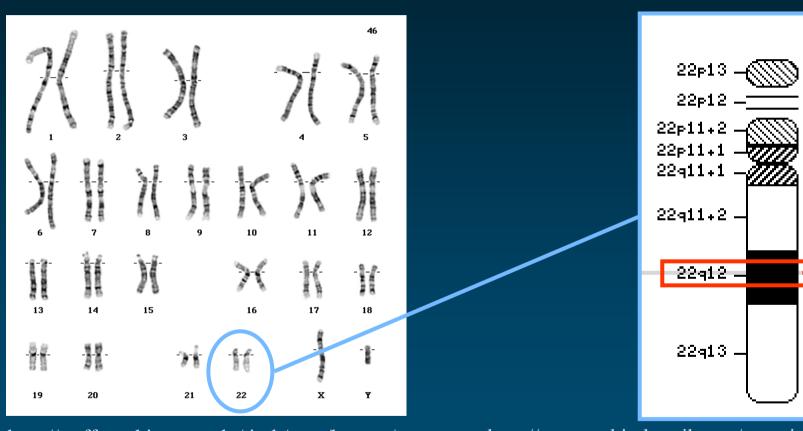


http://www.mayoclinic.com





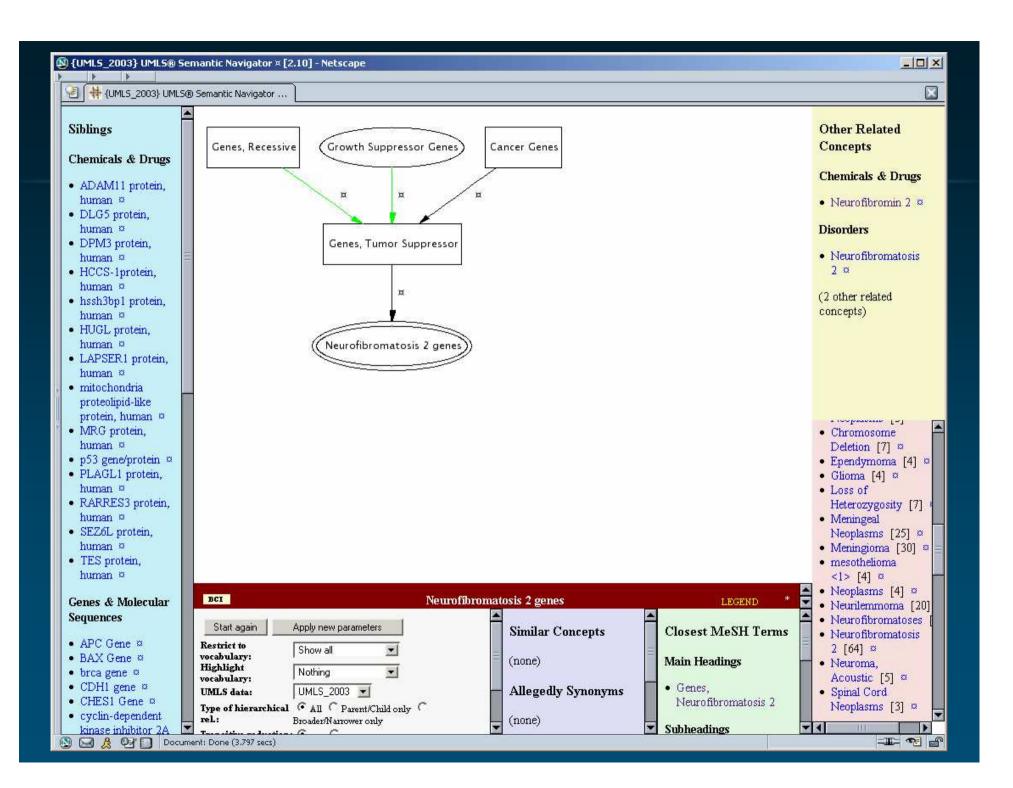
NF2 gene



http://staff.washington.edu/timk/cyto/human/



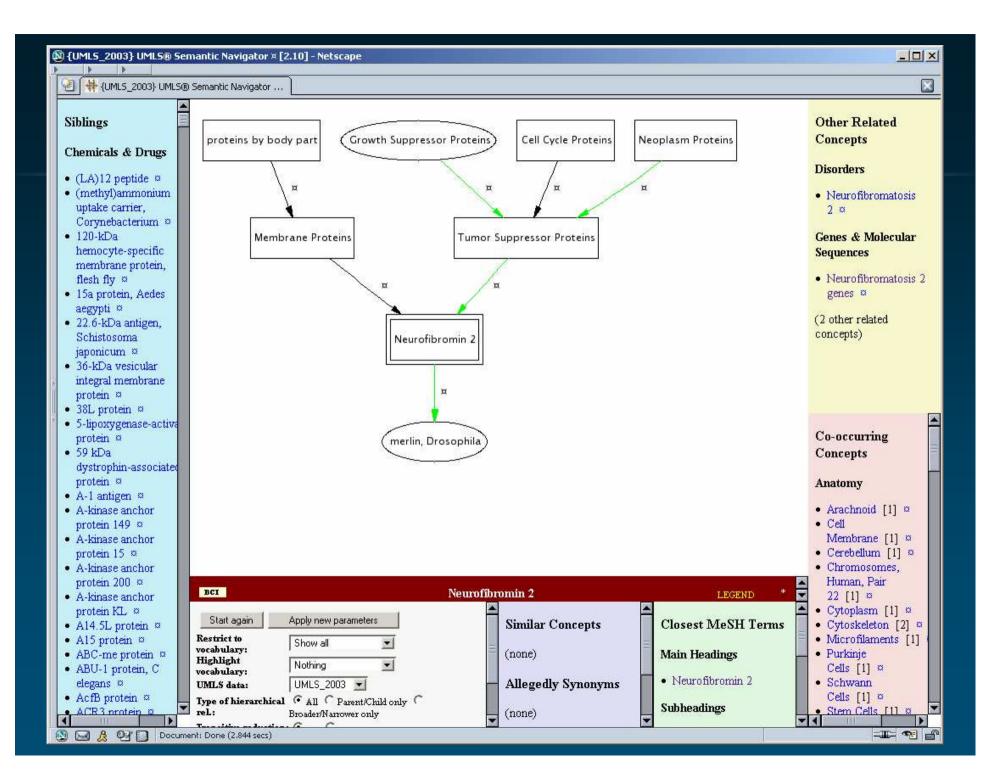


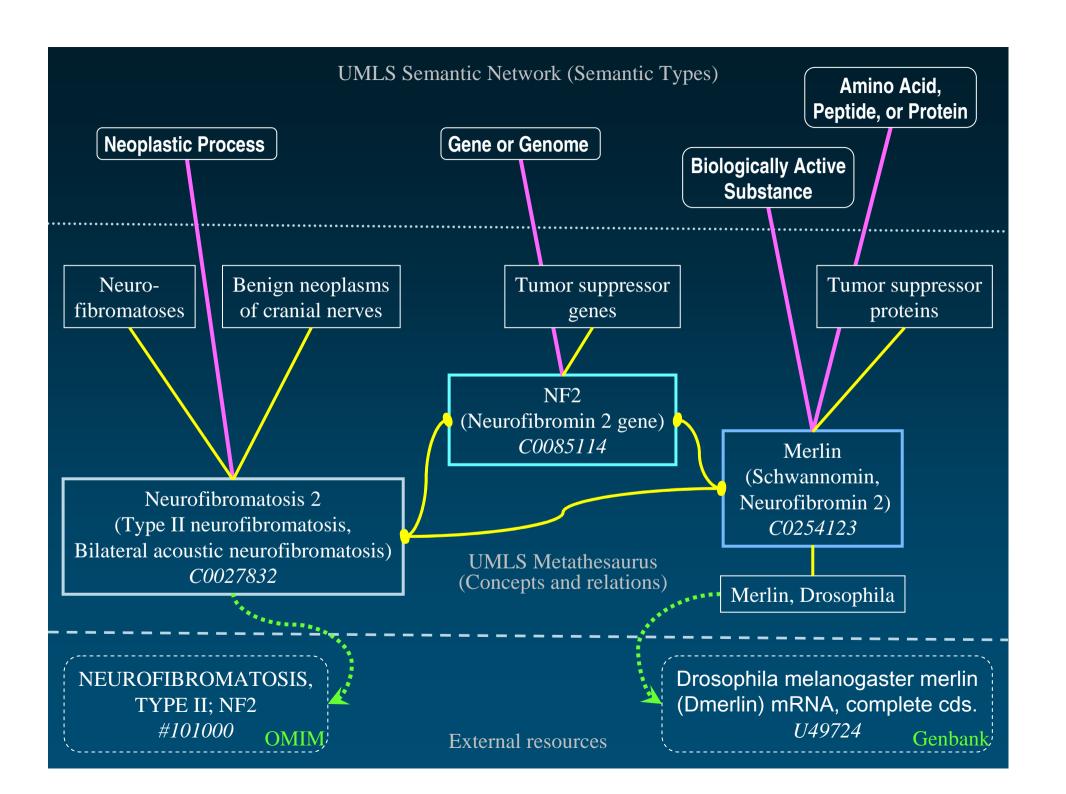


Merlin

- Synonyms
 - Neurofibromin 2
 - Schwannomin
 - Schwannomerlin
 - Neurofibromatosis-2
- ◆ 10 isoforms
- **♦** Annotations
 - Negative regulation of cell proliferation
 - Cytoskeleton
 - Plasma membrane







Limitations

- Genes not systematically represented
 - Most gene products and diseases are
- ◆ Gene/Gene product-Disease relations
 - Not systematically represented
 - Not explicitly represented (e.g., co-occurrence)
- Cross-references not systematically represented
- ◆ Naming conventions (genes)



Applications (1)

 $\overline{GenesTrace^{TM}}$

I.N. Sarkar & al. Columbia University

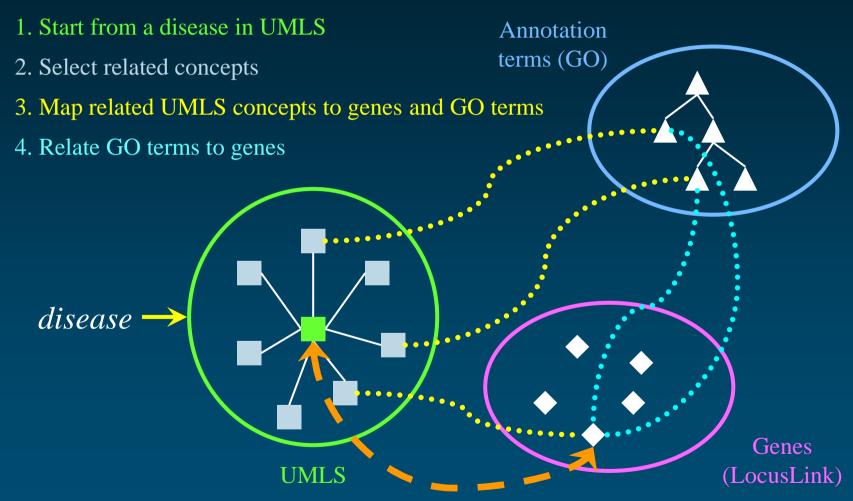
Objectives

◆ Relate diseases to genes through structured, integrated terminologies

◆ Biological Knowledge Discovery

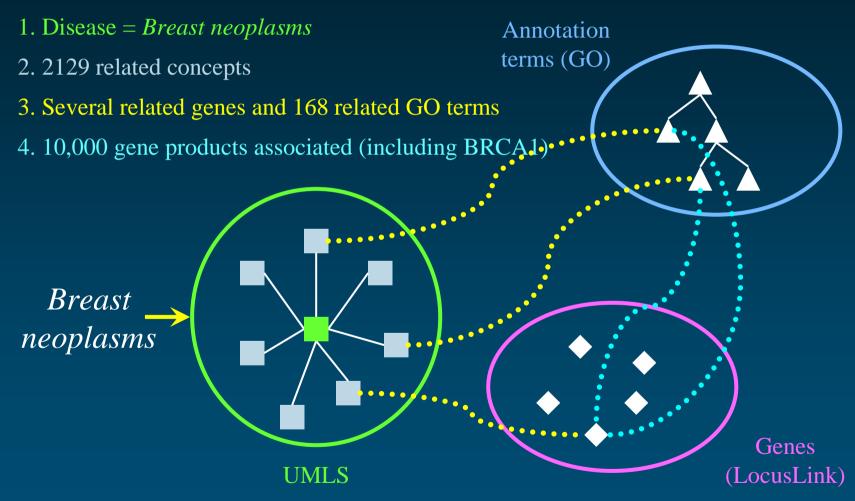


Resources and Methods





Validation Breast cancer – BRCA1 association





Limitations

- ◆ Noise
 - Too many non-specific GO terms associated (e.g., *nucleus*)
 - Too many genes associated
- ◆ But
 - Promising preliminary results
 - Room for refinement



References

◆ I. Sarkar, M. Cantor, O. Bodenreider, Y. Lussier. GenesTraceTM: Biological knowledge discovery via structured terminology – A feasibility study. Submitted to MEDINFO 2004.



Applications (2)

BioMeKe

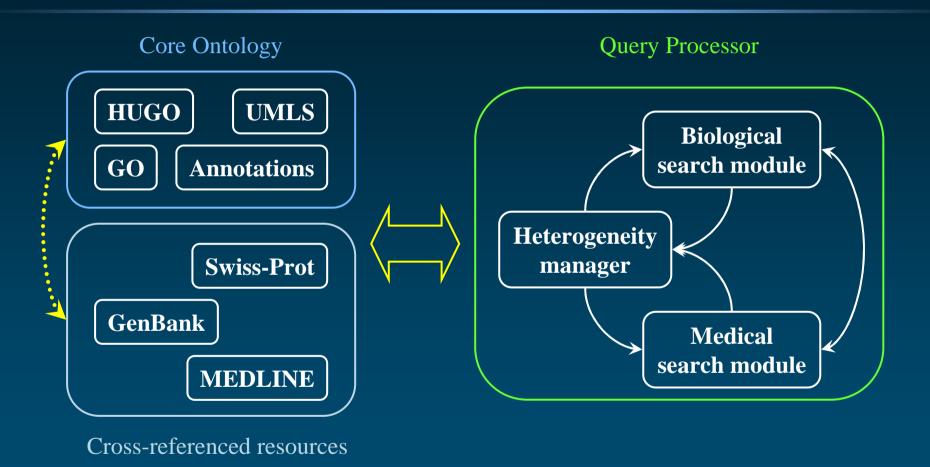
G. Marquet & al. LIM, Univ. Rennes, France

Objectives

- ◆ To develop a knowledge warehouse for transcriptome analysis (liver diseases)
- ◆ Semantic interoperability
 - Medical knowledge bases
 - Molecular biology and genetics knowledge bases



Components



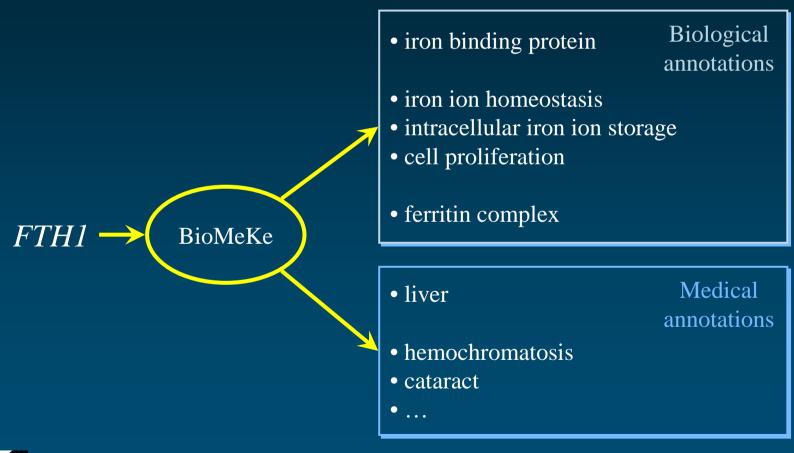


Example

- ◆ Input: *ferritin, heavy polypedpide 1*
- Mapping to biological resources
 - Not found in the Core ontology
 - Official name Ferritin heavy chain found through Xref
- Biological information obtained from GOA
- Mapping to medical resources
 - Not found in UMLS
 - Synonym *Ferritin H* found through Xref (Swiss-Prot)
- ◆ Medical information obtained through cooccurrence of MeSH index terms in MEDLINE



Results





Limitations

- Non-formal ontologies
 - Knowledge may be inconsistently represented
 - Knowledge may be implicit (mappings)
- ◆ Partial automation
 - User input required to select databanks, reformulate queries
- Semantic integration
 - Naming issues
 - Mappings must be updated regularly



References

◆ G. Marquet, C. Golbreich, A. Burgun. From an ontology-based search engine towards a more flexible integration for medical and biological information. Semantic Integration Workshop, ISWC 2003, Sanibel, Florida, October 20, 2003;61-67.



Conclusions

Conclusions

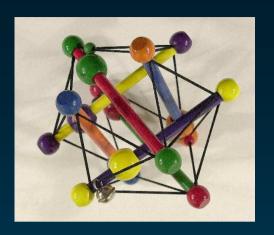
- ◆ Terminology integration provides some degree of information integration
- Most terminologies and the cross-referenced databases are readily available
- ◆ Lack of consistent representation



References

- ♦ UMLS umlsinfo.nlm.nih.gov
- **◆** UMLS browser
 - Knowledge Source Server: umlsks.nlm.nih.gov
 - Semantic Navigator: KSS, under UMLSKS resources
 - (free, but UMLS license required)
- UMLS and information integration
 - O. Bodenreider. The UMLS: Integrating biomedical terminology. *Nucl. Acids Res.* 2004;32(1) (in press)





Medical Ontology Research

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